

>SEQID#1_SGP037

ATGTCAACAGCTGCCCTTAATTACTTTGGTCAGAAAGTGTGGGAACCCAGGTGAGAAAGGAGAGTGTCTGCTAAGCTCCCGCCTGCTGCAGGACGACA
GGCGGTGACACCCACGTCGCCACAGCTCCACTTCAGAGCCTAGGTGTCTCGGTTTGACCCAGATGGTAGTGGGAGTCCAGCTACCTGGGACAA
TTTGGGATCTGGGATAACCGCATTTGATGAGCCAAATTCGTGCCACCCAGCAATTAAGTATGGCAAGCCAAATTCGCCAAATCAAGCTTGGAAATG
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TGATGGACAAGGTGGACCTGCAGCAGCTGATTTCTGTCTACATCCCAATGGAGAAATGTATTAATGGATTTGCTTCTTAAGGAGAAATCACTTGGAA
ACTCTGTGACCTTGGCTTTTCTAGAAATAGATAAAGCCTTTTCGAGTCAATGCCCGCCTGTCTGTGATGCAACTCTTCTGACCTCTGGGACTACT
GCAACAGTAGCCCTATJGCGAGATGGTATTGAACCTGGTTGAGCCAGTGTGGGACAGCCGGGCTATTTGTGTAGAAAGGAAACCCATGA
AGCTGACCAATTGACGATCTCCAGAAAGAAAGATGAAGAAAGGATCAAGAAATGTGTGGTTTGTAGCTTGGAAATAGTTTGGGCGAGC
CTCACGTAAATGGCAGGCTTGCAATGACAAAGATATTGGAGATTGGACCTTAAGACAGTGTGTGTCATAGCAGAACCTGAACCTAAGAGGAT
TAAGTTACATCATGCTGATGACAGCTTCTGCTCCTCACACAGATGGAAATTAACCTCATGTGTGAATAGTCAAGAGATTTGTGACTTTGTCAATC
AGTGCCATGATCCCAACGAAGCAGCCCATGGCGTGACTGAACAGGCAATAACAGTACTGAGGATAACAGTACTGCAGTAGTAGTGCCTTT
TGGTGCCTGGGGAAAATATAAGAACTCTGAAATCAACTTCTCATTTCAGCAGAAAGCTTTGCCCTCCAGTGACGATGGGCTGA

Fig. 1



>SEQID#2_SGP037
MSTAALITLVRSGGNQVRRRVLLSSRLQDDRRVPTTCHSSTSEPRCSRFDDGSGSPATWDNFGIWDNRDEPILLPSIKYKPKPKISLENVGCASQIG
KRKENEDREFDFAQLTDEVLYTAVYDGHGGPAAADFCHTHMEKCMDLLPKEKNLETLTLAFLIDKAFSSHARLSADATLLTSGTTATVALLRDGIEL
VVASVGDSRAILCRKGKPMKLTIDHTPERKDEKERIKKCGGFVAWNSLGQPHVNGRLAMTRSIGDLDLKTSGVIAEPETKRUKLHHADDSFLVLTIDGI
NFMVNSQHEICDFVNQCHDPNEAAHA VTEQAIQYGTEDNSTAVVPFGAWGKYKNSEINFSSRSFASSGRWA

Fig. 2